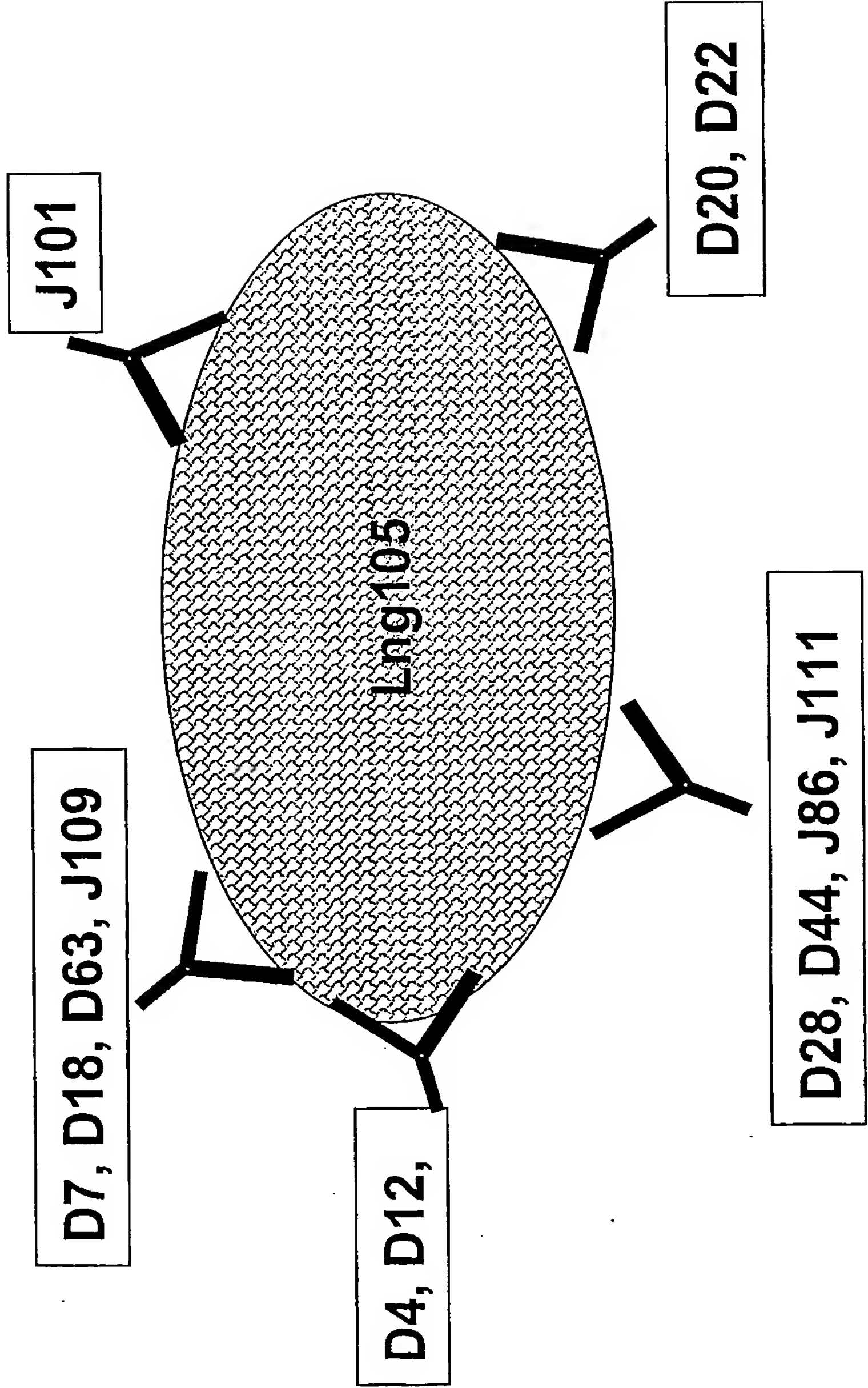


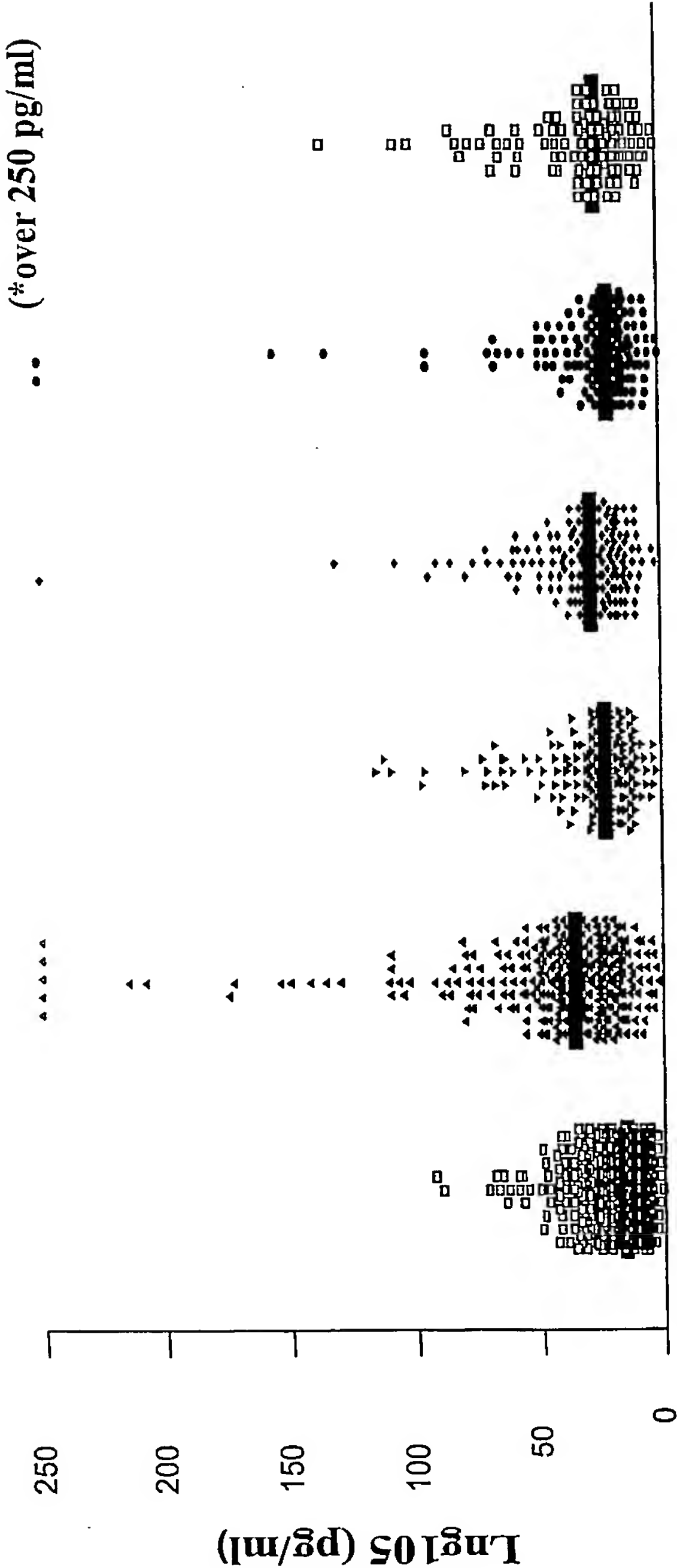
1/10

Figure 1



2/10

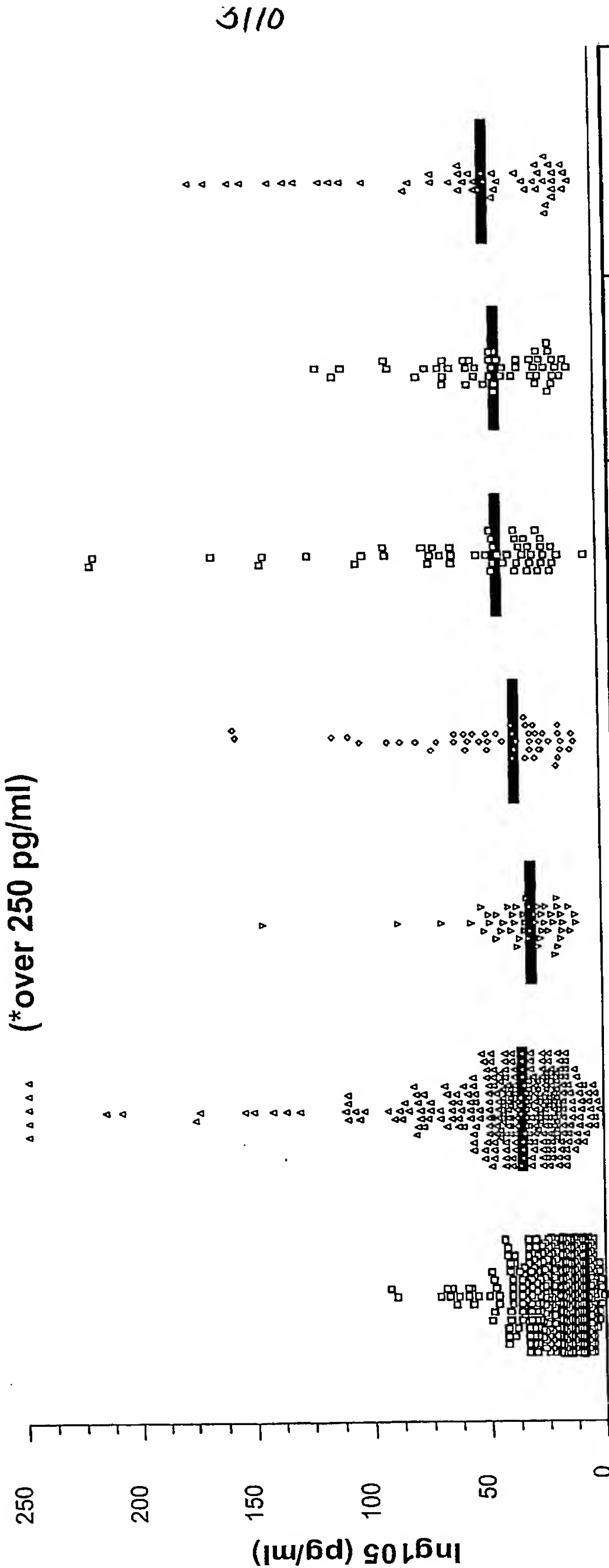
Figure 2



	normal	Lung Cancer	Prostate Cancer	Breast Cancer	Ovarian Cancer	Colon Cancer
N	555	296	163	158	151	100
Median	16.1	36.2	23.1	28.4	20.8	24.9

Anti-Lng105 MAbs D18/D28

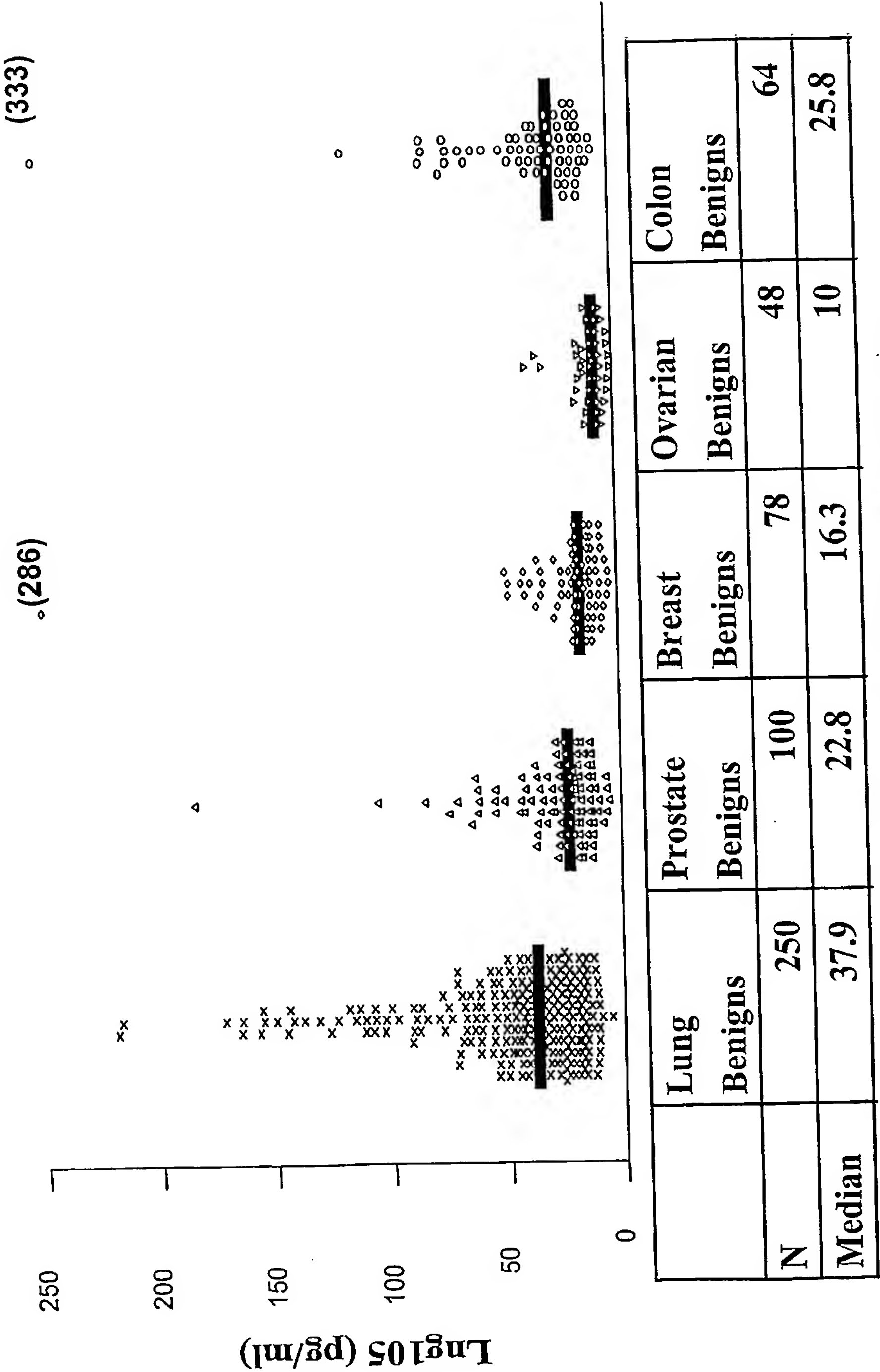
Figure 3



Anti-Lng105 MAbs D18/D28

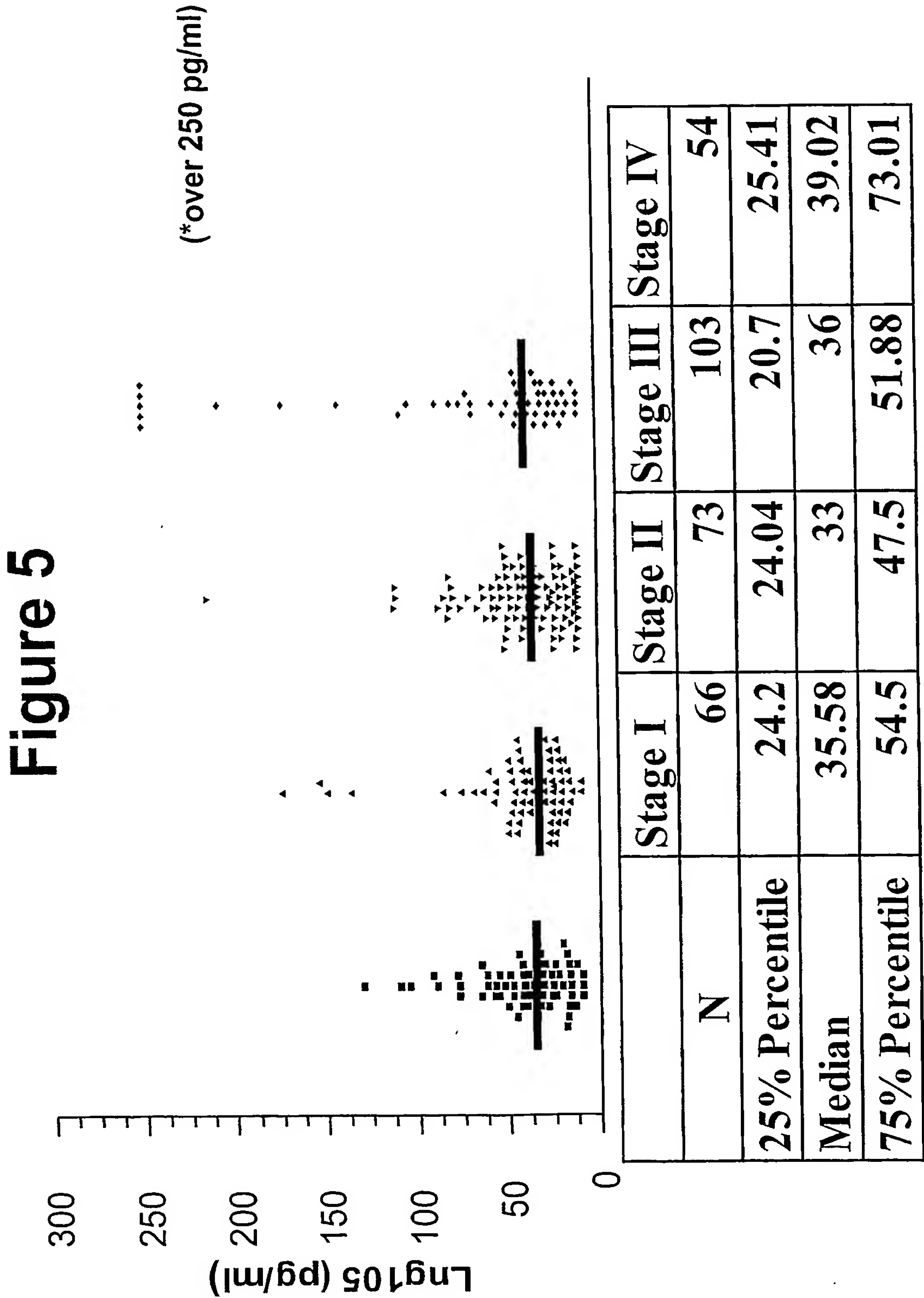
4/10

Figure 4



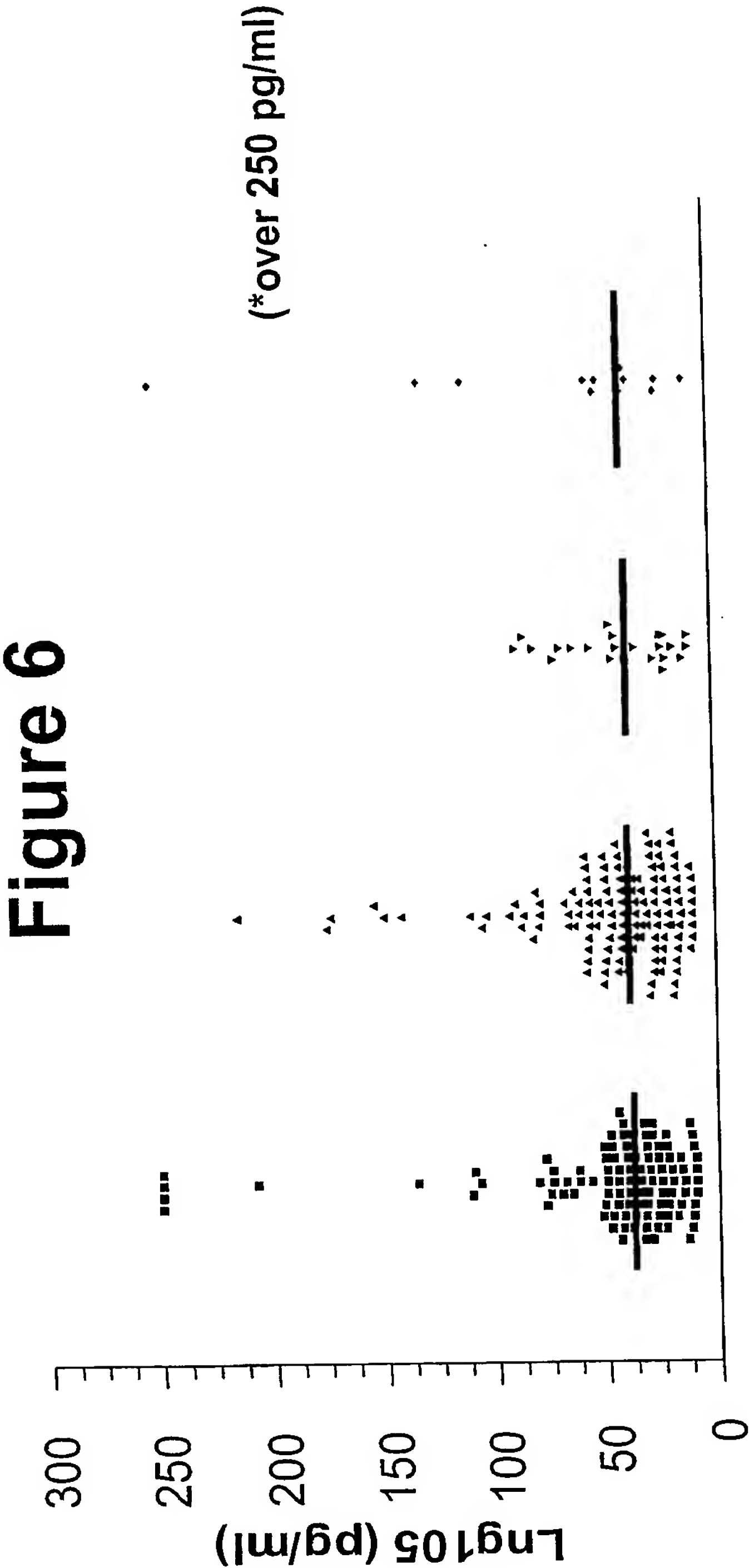
Anti-Lng105 MAbs D18/D28

5/10



Anti-Lng105 MAbs D18/D28

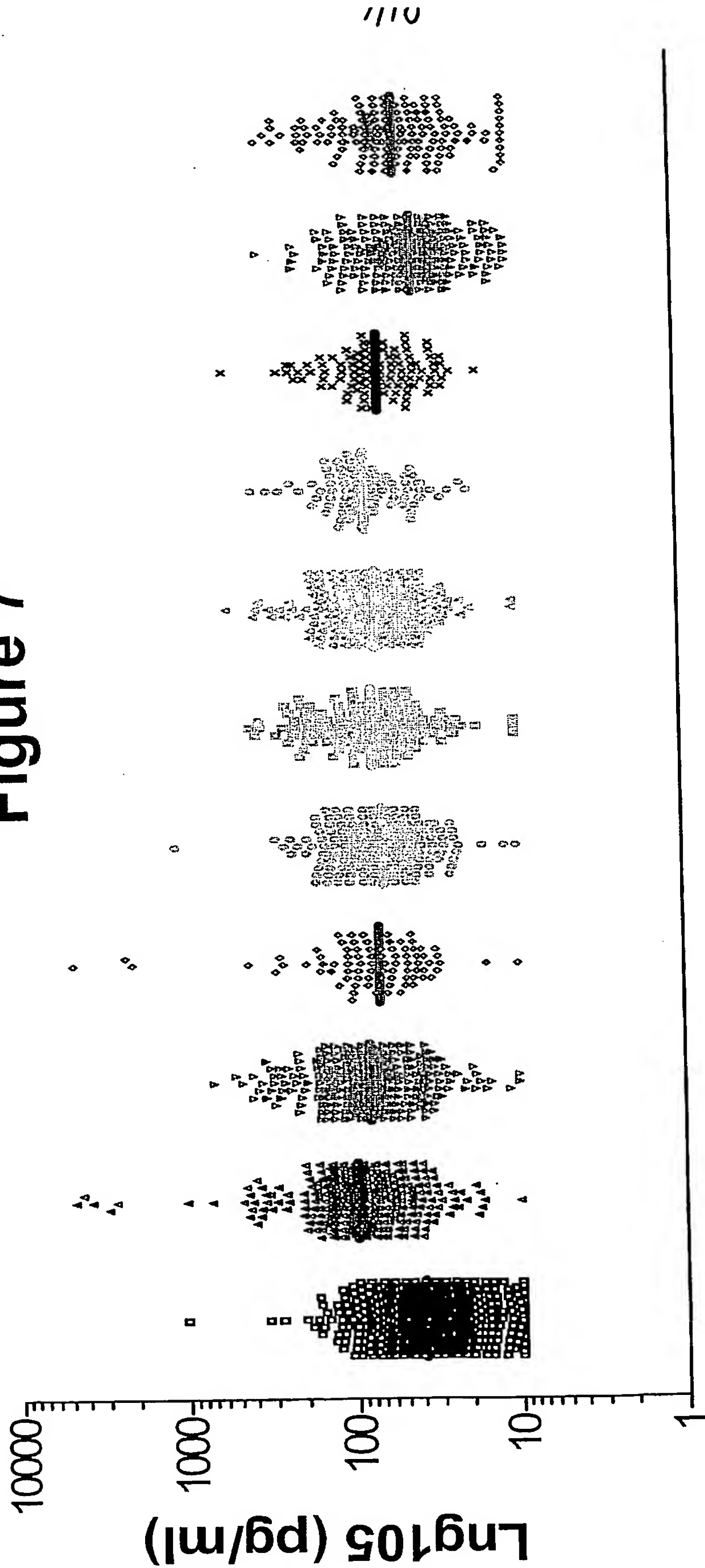
6/10



	Adeno-carcinoma	Squamous cell carcinoma	small cell carcinoma	large cell carcinoma
N	106	131	23	14
Median	38.0	38.8	38.1	39.1

Anti-Lng105 MAbs D18/D28

Figure 7

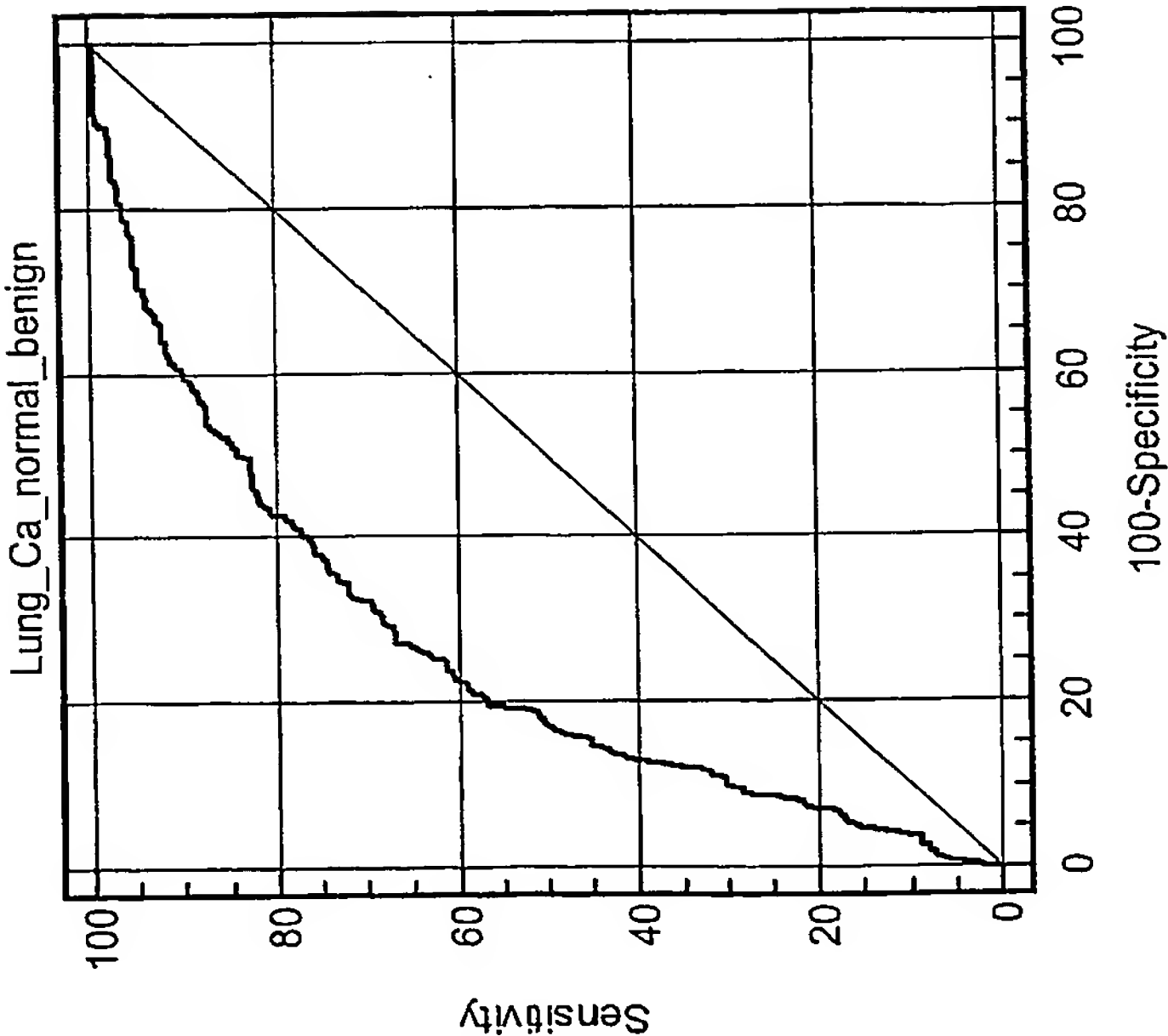


X Labels	Normal	LCa	L benign	BCa	B benign	CCa	C benign	PCa	P benign	OCa	O benign
Number of values	554	298	250	85	180	124	200	80	100	210	150
Minimum	10.00	10.00	10.00	10.00	10.00	10.00	10.00	18.31	15.64	10.00	10.00
25% Percentile	25.19	60.79	49.00	47.52	44.76	44.57	45.91	47.09	40.25	23.69	26.82
Median	39.28	99.20	83.11	71.10	65.88	74.32	68.90	77.58	62.56	37.73	47.22
75% Percentile	60.27	140.9	135.1	107.3	103.2	129.9	101.6	102.8	85.41	60.16	72.12
Maximum	1030	4710	689.4	4693	1121	391.6	524.1	355.1	520.5	310.4	311.3

Anti-Lng105 MAbs J109/J86

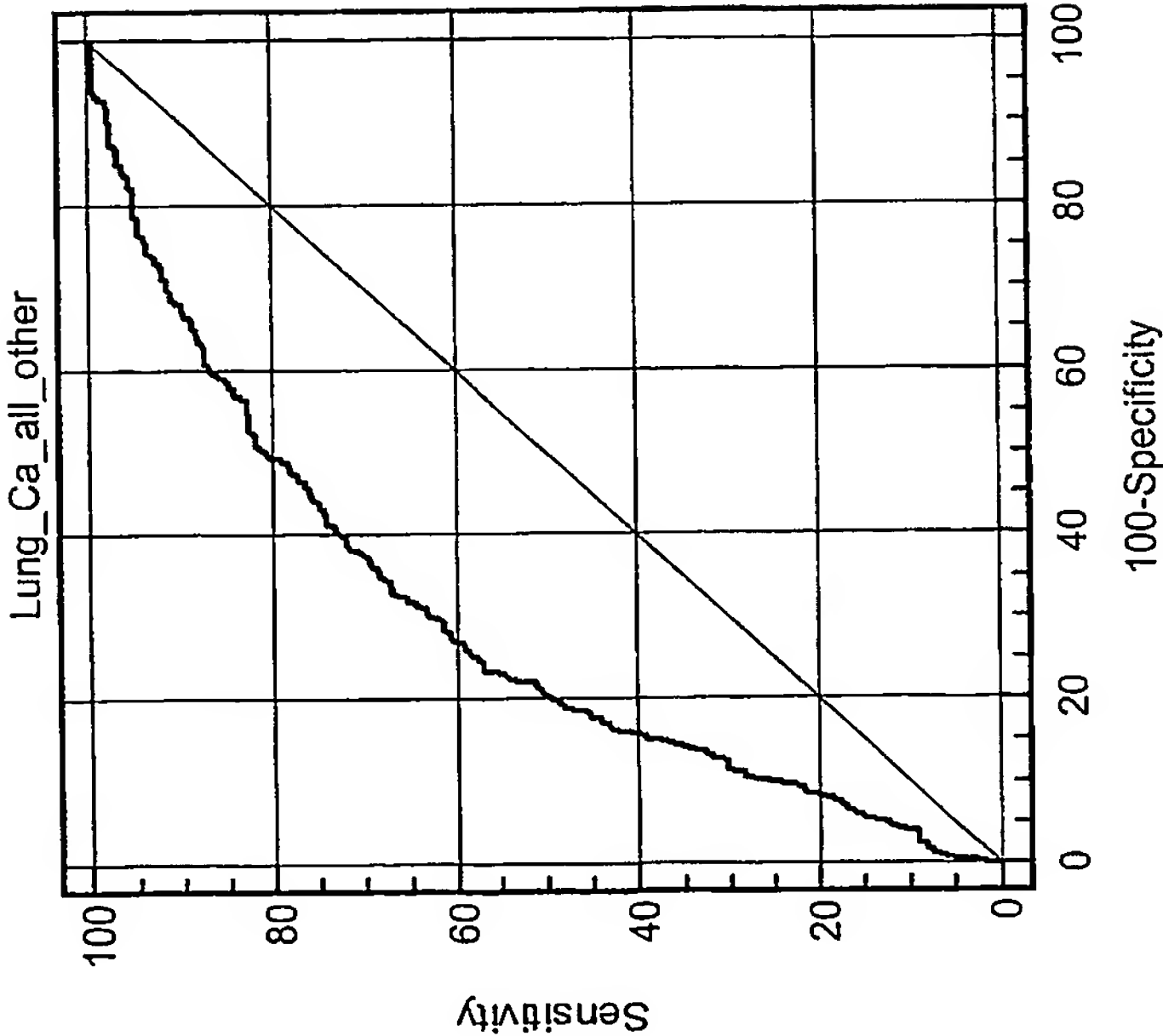
8/10

Figure 8



Lung Cancer Samples = 298
Normal and benign disease Samples = 804
Area under the ROC curve = 0.747
Standard error = 0.018
95% Confidence interval = 0.720 to 0.773

Figure 9



Lung Cancer Samples = 298
Normal, benign disease and non lung cancer
Samples = 1933
Area under the ROC curve = 0.714
Standard error = 0.018
95% Confidence interval = 0.694 to 0.732

Figure 10

